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When ligands and receptors are both attached on surfaces, because of the restriction of configurational freedom, their recognition kinetics may be substantially reduced as compared with freely diffusing species. In nature, this reduction may influence the efficiency of the capture and adhesion of circulating cells. Here we show that similar consequences are observed for colloids grafted with biomolecules that are used as probes for diagnostics. We exploit Brownian magnetic colloids that self-assemble into linear chains to show also that the resulting one-dimensional confinement considerably accelerates the recognition rate between grafted receptors and their ligands. We propose that because confinement significantly augments the colliding frequency, it also causes a large increase in the attempt frequency of the recognition. This work gives the basis of a rapid, homogeneous, and highly sensitive bioanalysis method.

association rate | bioassay | magnetic colloids | specific adhesion

Diagnostics techniques are generally based on building a specific immunocomplex structure in which the antigen to be detected is recognized by two antibodies (1–8). Early immunochemistry was based on precipitation of large complexes made of antibodies and antigens. Following the same track, the use of Brownian particles significantly improved the detection sensitivity, because of the increase in scattered light when aggregation between grafted colloids takes place. As the most simple but very generic example, let us consider an antigen *Ag* having two different epitopes for two antibodies, *A* and *B*. To reveal the presence of such antigen, particles grafted with *A* and *B* antibodies are mixed with the sample to be analyzed. The formation of small clusters is then expected, at a rate depending on many factors. Change in light scattering due to the presence of these small clusters will reveal the existence of sandwich-like structures: *A–Ag–B* (latex agglutination immunoassay). These homogeneous assays, as opposed to heterogeneous assays in which washing steps are necessary before detection, are today by far the most simple and straightforward assays. They were introduced >40 years ago (2), and today several hundred different tests based on this principle can be found on the market, mainly for infectious disease detection and protein quantification, as long as the antigen concentration to be detected is ≥ 1 nmol/liter (1). Meanwhile, many diagnostic assays require sensitivity in the picomolar range. They are presently performed by using a heterogeneous approach (such as ELISA) (1). For that range of concentration, if we were still to consider the same homogeneous approach, it would in principle take too long. Indeed, decreasing the number of antigens to be detected implies also decreasing the number of particles in solution,[†] and thus the particles' colliding frequency. Therefore, at these very low antigen concentrations, the encounter frequency between species becomes a critical issue to consider.

In this work, we demonstrate conditions that accelerate specific recognition between grafted ligands and receptors, giving the basis of a rapid, homogeneous, and highly sensitive bioanalysis method. Under a homogeneous magnetic field, Brownian magnetic colloids can transport and hold the reactants in the

vicinity, considerably increasing the colliding frequency. Because the used particles are superparamagnetic with a high susceptibility, the resulting magnetic colloidal forces induce a fast chaining process (9): the time scale for bringing two colloidal particles at contact in the presence of a magnetic field *H* and at an initial volume fraction ϕ , is given by

$$\tau = \frac{6\eta}{\phi\mu_0\chi^2 H^2}, \quad [1]$$

where η is the viscosity of the surrounding fluid, μ_0 is the vacuum magnetic permeability, and χ is the magnetic susceptibility of the particles (10). For typical experiments ($B = \mu_0 H = 20$ mT, $\chi = 0.95$, $\phi = 0.03\%$, and $\eta = 0.001$ Pa·s), the time τ to nucleate chains is < 1 s. These chains persist as long as the field is maintained and allows for rapid formation of ligands–receptors–ligands links between pairs of particles within the chain. To quantify the influence of this one-dimensional confinement on the recognition rate, we detect the resultant colloidal doublets that remain after the field is switched off, as illustrated in Fig. 1.

As a quantitative demonstration, we use ovalbumin as a model ligand. Colloidal magnetic particles 200 nm in diameter (Ademtech, Pessac, France) are grafted with polyclonal IgG rabbit anti-ovalbumin antibodies as a model for receptors (Bertin Technologies, Montigny-le-Bretonneux, France), with ≈ 30 antibodies per particle. The colloids volume fraction ϕ is 0.03% and corresponds to a particle concentration C_p of 120 pmol/liter. A final concentration C_{ova} of ovalbumin is adjusted, each sample is first incubated for 1 min at 25°C, and then a homogeneous field of 20 mT is applied for 5 min. The same experiment is also performed with zero field. Optical density, defined as $OD = -\log(I/I_0)$, where *I* and I_0 are, respectively, the transmitted and incident light intensity, is measured at a wavelength of 700 nm before and after the field is applied. The difference is plotted in Fig. 2 as a function of C_{ova} . In the absence of field, the signal is low, reflecting the very limited number of persisting doublets and therefore the inefficiency of free Brownian collisions on that time scale. By contrast, in the presence of a 20-mT field, a turbidity difference can be measured down to picomolar concentrations. For C_{ova} much less than C_p , only doublets of particles can form, allowing a straightforward determination of the absolute concentration of links C_{link} from this turbidity difference (see *Methods*). The slope of C_{link} as a function of C_{ova} in the presence of 20-mT magnetic field is found to be close to 1 (Fig. 3), which demonstrates that essentially all of the antigens are

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[†]In the regime of low antigen concentration, the particle number is large compared with the antigen number, so only doublets of particles can form. The light scattered by nonaggregated beads will add to the signal originating from the doublets. Turbidity measurements typically resolve better than a 1% intensity variation, so the concentration of beads in solution must not exceed 100 times the concentration of antigen.

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capture of the antigen on the beads. For free antigen in solution ($R_B = 5$ nm, $r_p = 0.25$ nm) and homogeneously reactive spheres ($R_A = 100$ nm, concentration 10^{-10} M), $k = 3 \times 10^{10}$ M⁻¹·s⁻¹, and $f = 0.2$, which leads to τ_1 of ≈ 2 s. The second step consists of bridging one colloidal particle, which has already captured one antigen, with another particle. In the absence of field, only free diffusion is involved, and τ_2 is estimated to be ≈ 1 min (homogeneously reactive particles at concentration 10^{-10} M with radius $R_B = 100$ nm and $r_p = 0.25$ nm, so $k = 6 \times 10^9$ M⁻¹·s⁻¹ and $f = 3 \times 10^{-2}$). Note that these two predicted time scales are certainly underestimated and should be considered as upper limits, because the assumption of uniformly reactive spheres is too optimistic. We have indeed measured 30 antibodies per particle, which is far below the estimated monolayer threshold of $\approx 1,000$ per particle (‡, 13).

In our experiment, the first step requires an incubation time of < 1 min, in good agreement with the previous diffusion-based estimation. By contrast, in the absence of field, the second step would require > 8 h, as deduced from the slope of Fig. 3. We can therefore conclude that the previous diffusion-based model does not hold once grafted biomolecules are concerned. Some repulsive colloidal forces could be responsible for this slowdown; however, because we do not find any role of the field strength above the onset of chaining, we can rule out this contribution as a major effect. We believe that the immobilization of the biomolecules onto colloidal surfaces causes restrictions to their dynamics, which drops further down the steric factor as compared with a nongrafted case. To improve the rate of colloidal bridging, we must therefore augment the attempt frequency. One simple way consists of increasing the particles' colliding frequency by locally increasing the particle concentration; this result is readily obtained from chaining magnetic colloids with magnetic field. As deduced from our experiment, the measured

agglutination rate is increased by a factor of 100 above the onset of chaining. This effect shifts the detection threshold of the latex agglutination immunoassay homogeneous bioassay down to 1 pmol/liter, making these simple methods as sensitive as heterogeneous approaches. However, although confinement has significantly increased the agglutination rate, it also raises the delicate question of the coupling between translation and rotation in such confined geometry and therefore the ultimate limitation of this approach.

This letter gives the basis of a previously unreported homogeneous assay for protein detection with high sensitivity. Indeed, because magnetic colloids can immediately self assemble under field, they are very efficient carriers, considerably reducing the diffusion time; most importantly, magnetic forces can hold colloids and reactants in vicinity, which increases the sampling frequency, allowing a more rapid reaction. These results also demonstrate the relatively slower process of recognition when both ligand and receptor are grafted onto colloidal surfaces.

Methods

Turbidity measurement relies on the fact that a doublet scatters more light than two separate beads. If α is the ratio of the total scattered light intensity by one doublet to the total scattered light intensity by one particle, the optical density difference before and after the magnetic field is applied will be $OD_a - OD_b = h\pi a^2 Q_{\text{scat}}(\alpha - 2)n_2/2.3$, where h is the optical path length, a is the particle radius, Q_{scat} is the scattering efficiency of one particle, and n_2 is the density number of doublets. $Q_{\text{scat}} = 0.23$ and $\alpha = 2.45$ are numerically computed by using Mie Theory (14) using freely available software (<ftp://ftp.eng.auburn.edu/pub/dmckwski/scatcodes/index.html>) with the measured particle optical index ($1.78 + 0.02i$ at 633 nm) and the measured particle diameter (200 nm).

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*Nevertheless, as shown by Berg and Purcell (13), a sphere partially covered by receptors behaves almost like a uniformly reactive sphere. Indeed, before diffusion moves the reactants out, they try many orientations during repeated encounters. The same argument explains why the steric factor f is higher than a purely geometrical estimation.

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